```
==============================================================
Sequence Listing could not be accepted due to errors.
See attached Validation Report.
If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).
Reviewer: Durreshwar Anjum
Timestamp: Fri Sep 14 12:53:23 EDT 2007
==============================================================



*************************************************************
Reviewer Comments:
116676-9.ST25.txt

1.  Delete this text which is on the headings as well as end of file.

2.  <140> US 10/597,787 <141> 2006-08-08
     <150> PCT/EP2005/001252 <151> 2005-02-08
      <150> DE 10 2004 006 249.8 <151> 2004-02-09
Use hard return enter and the format will be as below(please for the
similar errors).
<140>
<141>
<150>
<151>
3.  Seq Id 2:  Invalid response for  <213>, If <213> response is
Artificial,
     Unknown, please give the source of genetic material.



*************************************************************
```

Validated By CRFValidator v 1.0.3

Application No:     10597787          Version No:          1.0

Input Set:

Output Set:

Started:  2007-09-05 19:13:13.798
Finished:  2007-09-05 19:13:14.514
Elapsed:  0 hr(s) 0 min(s) 0 sec(s) 716 ms
Total Warnings:  4
Total Errors:  9
No. of SeqIDs Defined:  2
Actual SeqID Count:  2

| Error code | Error Description |
|------------|------------------|
| E  105 | Multiple identifiers on single line |
| E  105 | Multiple identifiers on single line |
| E  105 | Multiple identifiers on single line |
| E  248 | Order Sequence Error <150> -> <150>; Expected Mandatory Tag: <210> in Header |
| W  213 | Artificial or Unknown found in <213>  in SEQ ID (1) |
| W  213 | Artificial or Unknown found in <213>  in SEQ ID (2) |
| E  105 | Multiple identifiers on single line |
| E  257 | Invalid sequence data feature in <221>  in SEQ ID (2) |
| W  112 | Upper case found in data; Found at position(9) SEQID(2) |
| W  112 | Upper case found in data; Found at position(10) SEQID(2) |
| E  254 | The total number of bases conflicts with running total  Input: 0, Calculated : 17 SEQID(2) |
| E  254 | The total number of bases conflicts with running total, Input: 47, Calculated : 64 SEQID(2) |
| E  253 | The number of bases differs from <211>  Input: 47  Calculated:64 |

<110> SOMMERMEYER, Klaus

<120> PROCESS FOR THE PRODUCTION OF CONJUGATES FROM POLYSACCHARIDES AND POLYNUCLEOTIDES

<130> 116676-9

<140> US 10/597,787 <141> 2006-08-08

<150> PCT/EP2005/001252 <151> 2005-02-08

<150> DE 10 2004 006 249.8 <151> 2004-02-09

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<170> Patentln version 3.2

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<400> 1

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<210> 2

<211> 47

<212> DNA

<213> Artificial

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<221> misc_feature <222> (5)..(3)

<223> Spiegelmer

<400> 2

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